Table 1a: Characteristics of the study group consisting of 101 Turkish patients that underwent genetic analyses for CYP2C9*2 and *3, VKORC1 -1639 G>A and factor VII -401 G>T polymorphisms.

| | n (%) |
|---|----------------------------|
| Age in years | 63.74 ± 11.43 |
| (incan+SD) Daily warfarin dose | $4.07 \pm 1.60(1.13-7.86)$ |
| in mg, mean (range) | 1.07 - 1.00 (1.15 7.00) |
| Gender | |
| Female | 60 (59.41%) |
| Male | 41 (40.59%) |
| Population | 101 (100%) Turkish |
| Indication for warfarin treatment ^a | |
| AF | 68 (67.33%) |
| MVR | 21 (20.79%) |
| AVR | 9 (8.91%) |
| DVT | 5 (4.95%) |
| CAD | 21 (20.79%) |
| <i>CYP2C9*2</i> | |
| *1/*1 | 72 (71.3%) |
| *1/*2 | 23 (22.8%) |
| *2*2 | 6 (5.9%) |
| <i>CYP2C9*3</i> | |
| *1/*1 | 57 (56.4%) |
| *1/*3 | 34 (33.7%) |
| *3/*3 | 10 (9.9%) |
| VKORC1 -1639G>A | |
| GG (wild type) | 25 (24.8%) |
| GA (heterozygous) | 49 (48.5%) |
| AA (homozygous) | 27 (26.7%) |
| Factor VII -401G>T | |
| GG (wild type) | 47 (46.54%) |
| GT (heterozygous) | 37 (36.63%) |
| TT (homozygous) | 17 (16.83%) |

Data are presented as mean \pm standard deviation or number of patients (%), ^a: patients may have had more than one indication for warfarin therapy, DVT: deep vein thrombosis, MVR: mitral valve replacement, AVR: aortic valve replacement, AF: atrial fibrillation, CAD: coronary artery disease, *CYP2C9*: cytochrome P450 2C9 genotype, *VKORC1*: vitamin K epoxide reductase genotype, factor *VII: factor VII* genotype.